

0590
0530

#5

ENTERED



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/045,180A

DATE: 06/03/2002

TIME: 12:40:19

Input Set : A:\EP.txt

Output Set: N:\CRF3\06032002\J045180A.raw

3 <110> APPLICANT: Bougueleret, Lydie
4 Chumakov, Ilya
6 <120> TITLE OF INVENTION: Human Defensin Polypeptide Def-X, Genomic DNA and cDNA,
Composition
7 Containing Them and Applications to Diagnosis and to Therapeutic Treatment
9 <130> FILE REFERENCE: GEN-100D1
11 <140> CURRENT APPLICATION NUMBER: US 10/045,180A
12 <141> CURRENT FILING DATE: 2001-10-18
14 <150> PRIOR APPLICATION NUMBER: US 09/486,580
15 <151> PRIOR FILING DATE: 2000-02-25
17 <150> PRIOR APPLICATION NUMBER: PCT/FR98/01864
18 <151> PRIOR FILING DATE: 1998-08-28
20 <150> PRIOR APPLICATION NUMBER: FR 97/10823
21 <151> PRIOR FILING DATE: 1997-08-29
23 <160> NUMBER OF SEQ ID NOS: 12
25 <170> SOFTWARE: PatentIn version 3.1
27 <210> SEQ ID NO: 1
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29 <212> TYPE: DNA
30 <213> ORGANISM: Homo sapiens
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64 <222> LOCATION: (1111)..(1111)
65 <223> OTHER INFORMATION: n = a, c, g, or t.

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81 <221> NAME/KEY: TATA_signal
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89 <223> OTHER INFORMATION: n = a, c, g, or t.
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99 <221> NAME/KEY: misc_feature
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124 <222> LOCATION: (2155)..(2335)
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177 <221> NAME/KEY: misc_feature
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191 <223> OTHER INFORMATION: Exon 3
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195 <221> NAME/KEY: misc_feature
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197 <223> OTHER INFORMATION: Translation termination codon (TAA)
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202 <222> LOCATION: (4374)..(4379)
203 <223> OTHER INFORMATION:
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W--> 211 cctgtgcgta acaacatcac acnccaaatt taaccagctc tccccataac agcagcgctca      180
213 tgtgttactg aggaaatgcc tgtggattgg agtgtgttct gtgtgcagga ggtcgggtcca      240
215 ggtttcactt ctgcaggaca ctggacgttt cccaaaacca gcagactttc cccacgtgca      300

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| | | | | | | | |
|------|------------|-------------|-------------|-------------|-------------|-------------|------|
| 217 | cacacacccc | ttctcatttt | gcctctacat | ccatatccac | tgggccccttc | aggcacctac | 360 |
| 219 | taatgcctca | gaacctaaaa | coactcatctg | gggcccagtt | ccctgaatgg | ccctaattctc | 420 |
| 221 | ttctctgctg | ggaatgtagtc | ctctctccaac | cagtgcccac | cttgaaattg | ctgggctgct | 480 |
| 223 | acagatcagg | aactcactgc | ttctctcatag | gggcagccga | cttcactgct | ctgcaacagc | 540 |
| 225 | gaccacccct | agcgagcctt | gagatgcctc | ttgcctcctt | aagactgagg | gagacgtcttc | 600 |
| 227 | agctctcact | ccaactgcccc | aagtctctcca | cagcgcgggtg | ctgtctgcct | tcacacagag | 660 |
| W--> | 229 | ctgcaggggn | aggtcctgtg | tatccggcct | gctggaaccag | cgtgtgcac | 720 |
| | 231 | tggcaacagt | ggctgcccgg | cctgcacact | gggcttgcca | acctcctgct | 780 |
| | 233 | tcctccagga | gtgactgcat | tcttttccca | ttccagaaa | actgatgcca | 840 |
| | 235 | tatgaggagg | aggaggagga | ggagggttga | gagtggta | ttttaaattg | 900 |
| | 237 | tccttaggac | tcctccctca | ataaccocag | agggaccata | ccagctcatt | 960 |
| W--> | 239 | ccaagcatan | gagtaatcat | cccactcatg | ctgagtgtat | ggctggcatt | 1020 |
| | 241 | tgaactggct | ttagaacaag | gtgtttgagc | acacagcacc | gtcttctgctc | 1080 |
| W--> | 243 | ccctcccttg | tgagacctct | gagacacatt | naggtctcac | ctaaaaactc | 1140 |
| W--> | 245 | aggcccaaan | cggctctaaa | aaattgttca | gtctgaactc | tctaagggtca | 1200 |
| | 247 | tgtttgtctc | ctctaagaaa | ccacatgttg | ctgttatact | cttaattccg | 1260 |
| | 249 | caaacctgct | cctgcttagca | acacaagcgg | aggtggta | cctctcacc | 1320 |
| | 251 | caacacaccc | gtttgtccaa | acagctttga | tttgttttta | tagttggcag | 1380 |
| | 253 | aggagctggg | ttcaggccat | attccaaatc | ctcatctgtg | gtgtgattgc | 1440 |
| | 255 | tatctctctt | acagggttga | tactatgata | acagccocag | ctgtccagct | 1500 |
| | 257 | agtccttttg | tcacagatgt | tcagcctcag | ccaccagtga | ggatcaattg | 1560 |
| | 259 | cttaggagtc | atggagaatt | catagttgtg | agctacctgg | gcttgccagc | 1620 |
| | 261 | agacaaggca | tcctctctgt | aactcctatt | ttaatgccag | cttcccaaca | 1680 |
| | 263 | ctgctcttca | cagcaggtat | ttaaactact | caatagaaa | taacccctgaa | 1740 |
| W--> | 265 | cctgtttccca | aaagaccctt | aaatagggga | agtcctttcn | ctgtgtgtgc | 1800 |
| | 267 | atgtggcaac | atgaggcctg | ggacagggga | ctgtcctctg | cccactctgg | 1860 |
| | 269 | tagcttaaca | atctgtcagt | aatacaatac | aaaacttaaa | ctttcatact | 1920 |
| W--> | 271 | ccaggaaagt | gtgtttccaa | ctgacccctg | gattatgggg | ccacctcaga | 1980 |
| | 273 | tgagggaata | ttttgcaatc | tgggactgtt | ggttgctggg | ggcagtggtc | 2040 |
| | 275 | ttataaaact | caagcagttt | ccttccaaac | acacatgtcc | tacttaacgt | 2100 |
| W--> | 277 | gatgatcata | ctcatangct | gctaaaaacat | tanttttatt | ttgagaaaag | 2160 |
| W--> | 279 | ttcttgcccc | atggagtttt | cattnnatta | ntttatttat | tttcagaga | 2220 |
| | 281 | ctatgttgct | acagctggctc | tcacaactcct | gggctcaagg | ctatcttcta | 2280 |
| | 283 | tgaagagcgt | gagattgctc | gtgtgagcca | tctggggggc | tcactggccc | 2340 |
| W--> | 285 | cagattaatt | gttttttgc | attgaanttg | tttgacttcc | ttgtatatcc | 2400 |
| | 287 | ccattctaac | acgttagggtt | tgcataat | ttctctcatg | ttctgtgtgt | 2460 |
| | 289 | cagttgatgg | ttctcttgc | tgtgcaggtg | cttatgtgtt | caacgcagcc | 2520 |
| | 291 | attttccatt | tatttgctgt | tcctcttgat | gtcatagcca | agaaaataat | 2580 |
| | 293 | atgtccaaaa | gtctatctcc | tatatattct | tctagtattt | tatggtttca | 2640 |
| | 295 | ttaggtcttc | aatccattga | gttgattttt | gtatgttgta | taagaaaaaa | 2700 |
| | 297 | ataactctga | caaatctctaa | ggtagtatat | attagacaca | tacaattgct | 2760 |
| | 299 | acattgagct | gaaaaataa | aaacatattt | tatctttcaa | ccaactctat | 2820 |
| | 301 | ctgaacttgt | ttcacctata | gcctgatgag | gttgctgtcc | tctctacccc | 2880 |
| | 303 | ggagactgct | catccctcaa | cctcaaaaac | ccctctcatg | gggtgataat | 2940 |
| | 305 | cctgcacatg | attagttctc | tactacagtg | gaattcaggt | ctgttatagg | 3000 |
| | 307 | tcctgagaga | agagctctca | ttttcagaaa | ataagcagga | ttttatccgt | 3060 |
| | 309 | aattaaatca | ctgtttcgat | tactttttgc | aaatattaaa | gtaaaatttt | 3120 |
| | 311 | aaacagaaat | aatggttagg | tccttatcat | acccgtgaat | tccaagctag | 3180 |
| | 313 | aaacctagag | attcacacta | gaatgaaagc | tgggagagca | gaggagtctc | 3240 |

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315 ggagggccaat ggacacctgc aacctctcca acgaaatgcc tacctcctct cactgcagca 3300
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319 acatccactc ctgctctccc tctctctctc cagggtgacta cagttagag gccctccacc 3420
321 ctctctctct cttttctcct ggtggccctt caggcctggg cagagcgct ccaggccaaga 3480
323 gctcatgaga tgccagccca gaagcagcct ccagcagatg accagatgt ggtcattttac 3540
325 ttttcaggag atgacagctg ctctcttcag gtctcagggt agagatgcc gcatgcagag 3600
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329 cacttaggtg gctatactta acatctctgg tctctggaat tctcatatct aaatggaata 3720
331 gagaacccaa gaaatctaag agatttttct ttctccaaaa acttgattcc aagatatgac 3780
333 tgtgaaattc actagattta agatataagg agatgctacc tagttccttc tggagccaga 3840
335 caaacaaagt taagtatata ggaataatatt tcacctgtc tatataggag gttttagaac 3900
337 ctggagagga gccaaagaat gtgttcagggt gtgtgtgtga tgggcaggaa tgcagaaaaa 3960
339 tgaagcaaaq gagaatgagt ctgcaatcct gtgtgaccag cactgctctg tgtattttatt 4020
341 cctattgact gagattgttt gtgctaccgg ctgtaataca gccaacatca ctcatcagcc 4080
W--> 343 aacatgtgac ttctccaaga ttccctttac caccactgc tgnaccocgt actcagtttc 4140
345 tgatgctctc tctgggtccc caggtctaac aaagggtctg atctgceatt gcagagtact 4200
347 atactgcatt tttggagaac atcttggtgg gacctgtctc atcttggttg aacgtaccac 4260
349 aatctgctgc tactaagctt gcagactaga gcaaaaagagt tcaataattt ctttgagact 4320
351 taaagggaaat tgttattctt atacctgtgc ctgagatttc tgtctcctac ccaataataa 4380
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357 <211> LENGTH: 453
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359 <213> ORGANISM: Homo sapiens
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362 <221> NAME/KEY: CDS
363 <222> LOCATION: (52)..(336)
364 <223> OTHER INFORMATION: Def-X coding sequence
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369 Met Arg
370 1
372 acc ctc acc ctc ctc tet gcc ttt ctc ctg gtg gcc ctt cag gcc tgg 105
373 Thr Leu Thr Leu Leu Ser Ala Phe Leu Leu Val Ala Leu Gln Ala Trp
374 5 10 15
376 gca gag cag ctc cag gca aga gct cat gag atg cca gcc cag aag cag 153
377 Ala Glu Pro Leu Gln Ala Arg Ala His Glu Met Pro Ala Gln Lys Gln
378 20 25 30
380 cct cca gca gat gac cag gat gtg gtc att tac ttt tca gga gat gac 201
381 Pro Pro Ala Asp Asp Gln Asp Val Val Ile Tyr Phe Ser Gly Asp Asp
382 35 40 45 50
384 agc tgc tct ctt cag gtt cca ggc tca aca aag ggc ttg atc tgc cat 249
385 Ser Cys Ser Leu Gln Val Pro Gly Ser Thr Lys Gly Leu Ile Cys His
386 55 60 65
388 tgc aga gta cta tac tgc att ttt gga gaa cat ctt ggt ggg acc tgc 297
389 Cys Arg Val Leu Tyr Cys Ile Phe Gly Glu His Leu Gly Gly Thr Cys
390 70 75 80
392 ttc atc ctt ggt gaa cgc tac cca atc tgc tgc tac taa gcttcagac 346
393 Phe Ile Leu Gly Glu Arg Tyr Pro Ile Cys Cys Tyr

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RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:1; N Pos. 4123

VERIFICATION SUMMARY

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Input Set : A:\EP.txt

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L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:660
L:239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:960
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1080
L:245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1140
L:265 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1740
L:271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1920
L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:2100
L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:2160
L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:2340
L:343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:4080